

sub pg. 25

Peptide	Sequence
P12 ₍₃₂₂₋₃₃₃₎	PheCysLeuGlyProCysProTyrIleTrpSerLeuAspThr
P28 ₍₃₂₂₋₃₄₄₎	PheCysLeuGlyProCysProTyrIleTrpSerLeuAspThrGlnLysVal LeuAlaLeuTyr
P29 ₍₃₁₃₋₃₃₅₎	HisGluProLysGlyTyrHisAlaAsnPheCysLeuGlyProCysProTyr IleTrpSerLeuAspThr
P30	PheSerLeuGlyProCysProTyrIleTrpSerLeuAspThr
P31	PheCysLeuGlyProSerProTyrIleTrpSerLeuAspThr
P32	PheSerLeuGlyProSerProTyrIleTrpSerLeuAspThr
P33	PheCysLeuGlyProCysProTyrIleTrpSerAspAspAsp
P34	AspAspAspGlyProCysProTyrIleTrpSerLeuAspThr
P35	AspAspAspGlyProCysProTyrIleTrpSerAspAspAsp
P36	GlyProCysProTyrIleTrpSerAspAspAsp
P37	AspAspAspGlyProCysProTyrIleTrpSer
P38	AspGlyProCysProTyrIleTrpSerAsp

Fig. 6 shows the results of inhibition of TGF β 1 by the peptides in Table 3.

It can be seen from Fig. 6 that peptide P29 is active. This peptide includes the previously tested peptide P12 and has 9 extra amino acids towards the N-terminal end (Fig. 4). Investigations conducted by Quian SW et al. (1992) Proc. Natl. Acad. Sci. 89:6290-6294) and by Burmester JK et al. (1993) Proc. Natl. Acad. Sci. 90:8628-8632) using chimeric recombinant proteins identified a region of TGF β 1 that is necessary for the activity of this cytokine (amino acids 40 to 82 in the sequence of mature TGF β 1). It was speculated that peptide P29 (amino acids 34 to 56 in the sequence of mature TGF β 1), extending over a larger region than peptide P12 (amino acids 43 to 56), might acquire a three-dimensional structure more like the structure of the TGF β 1 in circulation. For this reason, peptide P29 was used for tests of binding to the cell receptors, based on affinity labelling.